



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/772,607C

DATE: 04/16/2003

TIME: 09:47:45

Input Set : A:\SEQLIST1.4-3-03.txt

Output Set: N:\CRF4\04162003\I772607C.raw

4 <110> APPLICANT: Jonassen, Ib
 5 Havelund, Svend
 6 Hansen, Per Hertz
 7 Kurtzhals, Peter
 8 Halstrom, John B.
 10 <120> TITLE OF INVENTION: Peptide Derivatives
 13 <130> FILE REFERENCE: 4409.214-US
 15 <140> CURRENT APPLICATION NUMBER: US 09/772,607C
 16 <141> CURRENT FILING DATE: 2001-01-30
 18 <150> PRIOR APPLICATION NUMBER: US 09/068,822
 19 <151> PRIOR FILING DATE: 1998-05-14
 21 <150> PRIOR APPLICATION NUMBER: PCT/DK96/00106
 22 <151> PRIOR FILING DATE: 1996-03-18
 24 <150> PRIOR APPLICATION NUMBER: DK 275/95
 25 <151> PRIOR FILING DATE: 1995-03-18
 27 <160> NUMBER OF SEQ ID NOS: 14
 29 <170> SOFTWARE: FastSEQ for Windows Version 4.0

**Does Not Comply
Corrected Diskette Needed**

ERRORED SEQUENCES

288 <210> SEQ ID NO: 14
 289 <211> LENGTH: 29
 290 <212> TYPE: PRT
 291 <213> ORGANISM: Artificial Sequence
 293 <220> FEATURE:
 294 <223> OTHER INFORMATION: Synthetic
 296 <220> FEATURE:
 297 <221> NAME/KEY: MOD_RES
 298 <222> LOCATION: LOCATION:29
 299 <223> OTHER INFORMATION: Lys at position 29 is modified with Nepsilon-gamma- Glu
 (Nalpha -
 300 tetradecanoyl)-OH)
 302 <400> SEQUENCE: 14
 303 Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly
 304 1 5 10 15
 305 Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr Lys
 306 20 25
 E--> 310 5

delete

VERIFICATION SUMMARY

DATE: 04/16/2003

PATENT APPLICATION: US/09/772,607C

TIME: 09:47:46

Input Set : A:\SEQLIST1.4-3-03.txt

Output Set: N:\CRF4\04162003\I772607C.raw

L:62 M:283 W: Missing Blank Line separator, <400> field identifier
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:310 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14